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RAW SEQUENCE LISTING   DATE: 05/01/2002
PATENT APPLICATION: US/09/930,020A   TIME: 09:21:25

          Input Set : A:\18501-31.app
          Output Set: N:\CRF3\05012002\I930020A.raw

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 3 <110> APPLICANT: Gish, Kurt C.
 4         Mack, David H.
 5         Wilson, Keith E.
 6         Eos Biotechnology, Inc.
 8 <120> TITLE OF INVENTION: Methods of Diagnosis of Colorectal Cancer, Compositions
 9         and Methods of Screening for Colorectal Cancer
10         Modulators
12 <130> FILE REFERENCE: 018501-003100US
14 <140> CURRENT APPLICATION NUMBER: US 09/930,020A
15 <141> CURRENT FILING DATE: 2001-08-14
17 <150> PRIOR APPLICATION NUMBER: US 09/663,733
18 <151> PRIOR FILING DATE: 2000-09-15
20 <160> NUMBER OF SEQ ID NOS: 3
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103 <223> OTHER INFORMATION: CBF9
105 <400> SEQUENCE: 2
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113       50                 55                 60
114 Gly Ser Phe Glu Arg Ser Lys His Phe Ala Ile Thr Val Cys Asp Gly
115 65                 70                 75                 80
116 Leu Asp Ile Ser Pro Glu Arg Val Arg Val Gly Ala Phe Gln Phe Ser
117                85                 90                 95
118 Ser Thr Pro His Leu Glu Phe Pro Leu Asp Ser Phe Ser Thr Gln Gln
119            100                105                110
120 Glu Val Lys Ala Arg Ile Lys Arg Met Val Phe Lys Gly Gly Arg Thr
121       115                120                125
122 Glu Thr Glu Leu Ala Leu Lys Tyr Leu Leu His Arg Gly Leu Pro Gly
123       130                135                140
124 Gly Arg Asn Ala Ser Val Pro Gln Ile Leu Ile Ile Val Thr Asp Gly
125 145                150                155                160
126 Lys Ser Gln Gly Asp Val Ala Leu Pro Ser Lys Gln Leu Lys Glu Arg
127            165                170                175
128 Gly Val Thr Val Phe Ala Val Gly Val Arg Phe Pro Arg Trp Glu Glu
129            180                185                190
130 Leu His Ala Leu Ala Ser Glu Pro Arg Gly Gln His Val Leu Leu Ala
131       195                200                205
132 Glu Gln Val Glu Asp Ala Thr Asn Gly Leu Phe Ser Thr Leu Ser Ser
133       210                215                220
134 Ser Ala Ile Cys Ser Ser Ala Thr Pro Asp Cys Arg Val Glu Ala His
135 225                230                235                240
136 Pro Cys Glu His Arg Thr Leu Glu Met Val Arg Glu Phe Ala Gly Asn
137            245                250                255
138 Ala Pro Cys Trp Arg Gly Ser Arg Arg Thr Leu Ala Val Leu Ala Ala
139            260                265                270
140 His Cys Pro Phe Tyr Ser Trp Lys Arg Val Phe Leu Thr His Pro Ala
141       275                280                285
142 Thr Cys Tyr Arg Thr Thr Cys Pro Gly Pro Cys Asp Ser Gln Pro Cys
143       290                295                300
144 Gln Asn Gly Gly Thr Cys Val Pro Glu Gly Leu Asp Gly Tyr Gln Cys
145 305                310                315                320
146 Leu Cys Pro Leu Ala Phe Gly Gly Glu Ala Asn Cys Ala Leu Lys Leu
147            325                330                335
148 Ser Leu Glu Cys Arg Val Asp Leu Leu Phe Leu Leu Asp Ser Ser Ala
149            340                345                350
150 Gly Thr Thr Leu Asp Gly Phe Leu Arg Ala Lys Val Phe Val Lys Arg
151       355                360                365
152 Phe Val Arg Ala Val Leu Ser Glu Asp Ser Arg Ala Arg Val Gly Val
153       370                375                380
154 Ala Thr Tyr Ser Arg Glu Leu Leu Val Ala Val Pro Val Gly Glu Tyr
155 385                390                395                400
156 Gln Asp Val Pro Asp Leu Val Trp Ser Leu Asp Gly Ile Pro Phe Arg
157            405                410                415
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160 Gly Phe Gly Ser Ala Thr Arg Thr Gly Gln Asp Arg Pro Arg Arg Val
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162 Val Val Leu Leu Thr Glu Ser His Ser Glu Asp Glu Val Ala Gly Pro
163         450             455             460
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165 465             470             475             480
166 Glu Ala Val Arg Ala Glu Leu Glu Glu Ile Thr Gly Ser Pro Lys His
167             485             490             495
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170 Leu Gln Gly Lys Leu Cys Ser Arg Gln Arg Pro Gly Cys Arg Thr Gln
171         515             520             525
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176 Phe Glu Val Asn Pro Asp Val Thr Gln Val Gly Leu Val Val Tyr Gly
177             565             570             575
178 Ser Gln Val Gln Thr Ala Phe Gly Leu Asp Thr Lys Pro Thr Arg Ala
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180 Ala Met Leu Arg Ala Ile Ser Gln Ala Pro Tyr Leu Gly Gly Val Gly
181         595             600             605
182 Ser Ala Gly Thr Ala Leu Leu His Ile Tyr Asp Lys Val Met Thr Val
183         610             615             620
184 Gln Arg Gly Ala Arg Pro Gly Val Pro Lys Ala Val Val Val Leu Thr
185 625             630             635             640
186 Gly Gly Arg Gly Ala Glu Asp Ala Ala Val Pro Ala Gln Lys Leu Arg
187             645             650             655
188 Asn Asn Gly Ile Ser Val Leu Val Val Gly Val Gly Pro Val Leu Ser
189             660             665             670
190 Glu Gly Leu Arg Arg Leu Ala Gly Pro Arg Asp Ser Leu Ile His Val
191         675             680             685
192 Ala Ala Tyr Ala Asp Leu Arg Tyr His Gln Asp Val Leu Ile Glu Trp
193         690             695             700
194 Leu Cys Gly Glu Ala Lys Gln Pro Val Asn Leu Cys Lys Pro Ser Pro
195 705             710             715             720
196 Cys Met Asn Glu Gly Ser Cys Val Leu Gln Asn Gly Ser Tyr Arg Cys
197             725             730             735
198 Lys Cys Arg Asp Gly Trp Glu Gly Pro His Cys Glu Asn Arg Glu Trp
199             740             745             750
200 Ser Ser Cys Ser Val Cys Val Ser Gln Gly Trp Ile Leu Glu Thr Pro
201         755             760             765
202 Leu Arg His Met Ala Pro Val Gln Glu Gly Ser Ser Arg Thr Pro Pro
203         770             775             780
204 Ser Asn Tyr Arg Glu Gly Leu Gly Thr Glu Met Val Pro Thr Phe Trp
205 785             790             795             800
206 Asn Val Cys Ala Pro Gly Pro
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Input Set : A:\18501-31.app
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211 <211> LENGTH: 5
212 <212> TYPE: PRT
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: Description of Artificial Sequence:conserved
217       cytokine receptor extracellular motif
219 <220> FEATURE:
220 <221> NAME/KEY: MOD_RES
221 <222> LOCATION: (3)
222 <223> OTHER INFORMATION: Xaa = any amino acid
224 <400> SEQUENCE: 3
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/01/2002
PATENT APPLICATION: US/09/930,020A   TIME: 09:21:26

Input Set : A:\18501-31.app
Output Set: N:\CRF3\05012002\I930020A.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 3

VERIFICATION SUMMARY   DATE: 05/01/2002
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Input Set : A:\18501-31.app
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L:225 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0